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TECH CENTER 1600/2900

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## RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/714,882

DATE: 05/29/2001

TIME: 11:46:13

Input Set : A:\Seqlist.txt

Output Set: C:\CRF3\05292001\I714882.raw

ENTERED

4 <110> APPLICANT: Turner, C. Alexander Jr.  
5 Nehls, Michael C.  
6 Friedrich, Glenn  
7 Zambrowicz, Brian  
8 Sands, Arthur T.  
10 <120> TITLE OF INVENTION: Novel Human Notch Ligand Proteins and Polynucleotides  
11 Encoding the Same  
13 <130> FILE REFERENCE: LEX-0091-USA  
C--> 15 <140> CURRENT APPLICATION NUMBER: US/09/714,882  
C--> 15 <141> CURRENT FILING DATE: 2000-11-16  
15 <150> PRIOR APPLICATION NUMBER: US 60/165,959  
16 <151> PRIOR FILING DATE: 1999-11-17  
18 <160> NUMBER OF SEQ ID NOS: 14  
20 <170> SOFTWARE: FastSEQ for Windows Version 4.0  
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23 <211> LENGTH: 2070  
24 <212> TYPE: DNA  
25 <213> ORGANISM: Homo sapiens  
27 <400> SEQUENCE: 1  
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29 atcaaagcag aggaacataa taaaagacaa aaggaaagaa atgtcaccac acaggtatca 120  
30 gtgaacgaaa tcaaacaata tttatcacac atatttgaac aaagaacatc tagtaatgta 180  
31 atcaataaaa gagaaaatct cctggagaaa aagaagaatc aacgtaaaat aagaataaaaa 240  
32 ggaattcaaa ataaagatat cttgaagaga aataagaatc atttacaaaa gcaagcagag 300  
33 aaaaatttta cagatgaagg agaccagcta tttaagatgg gcatcaagggt tctccagcag 360  
34 tctaaaagcc aaaaacaaaa agaagaagcc tacctacttt ttgccaaagc agctgacatg 420  
35 ggaaacttga aagctatgga gaaaatggct gacgctttgc tatttggaag ttttggcggtg 480  
36 caaaatataa cagcagctat ccaattatat gatgccttgg cttaaagaagg atcatgtaaa 540  
37 gcccaaaacg cattaggatt tttgtcttct tatggaatg gaatggaata tgatcaagct 600  
38 aaggcactga tatattacac ctttgggaagt gctggaggaa acatgatgtc ccagatgatt 660  
39 ttgggggtaca gatatttgtc gggaatcaat gttctacaga attgtgaagt tgccctaagt 720  
40 tattacaaga aagtggcaga ttatattgct gacacatttg aaaaaagtga aggtgttcca 780  
41 gtggaaaaag tgagactaac ggaaagacct gaaaatctga gttctaacag tgagattttg 840  
42 gattgggaca tataccaata ctataaattt ttggcagaaa gaggagatgt tcagatacaa 900  
43 gtctctcttg gacaattaca tctaattggc aggaaaggct tagatcagga ttactacaaa 960  
44 gcattacact acttcttaaa ggcagcaaag gccgggagtg caaatgccat ggcattttata 1020  
45 ggaaagatgt atttagaggg gaatgctgcc gtgccgcaa ataacgctac tgccttcaag 1080  
46 tacttttcca tggcagccag taagggaat gcaatcgcc ttcatgggct tggcttctct 1140  
47 tactttcatg gaaaaggagt tcccctgaat tatgccraag cacttaaata ctttcagaaa 1200  
48 gctgcggaag aagggtggcc cgacgcacag ttccagttag gcttcatgta ctactctggc 1260  
49 tctggaatat ggaaggatta taaacttgcc ttcaaattt tttacctggc atctcagagt 1320  
50 gggcagcccc tcgccattta ttatctggcc aagatgtatg caacaggaac aggagtagta 1380  
51 agatcatgca gaactgctgt ggagctttat aaagggtgtc gtgaactagg ccaactgggt 1440  
52 gagaaattcc tgacagctta ctttgccctat aaggatgggt atatagattc ttctcttgtt 1500  
53 cagtatgcac tgcttgacga aatggggtat gaagtagctc aaagcaattc agcattcatt 1560  
54 ttggaatcta aaaaggctaa cattcttgaa aaagagaaga tgtatccaat ggcgcttctc 1620  
55 ctatggaatc gagctgccat tcaaggcaat gcatttgcta gagtaaaaaat tggagattac 1680

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56 cattactatg gctatgggac taagaaagac tatcaaacag cagccacaca ctacagcatt 1740
57 gcagccaaca aataccacaa cgcgcaagcc atgttcaatc tggcttatat gtatgaacac 1800
58 ggcttaggca tcacaaagga cattcacttg gccagaagat tgtacgacat ggctgctcaa 1860
59 acgagtccag atgccacat acctgtgctc tttgccgtca tgaaactgga aactacgcat 1920
60 ttgctccggg atatcctgtt ttttaatcag ttcacaacga gatggaactg gctgaaactg 1980
61 gacaacacca ttggaccaca ctgggactta tttgtgattg gcctcattgt tcctgggctg 2040
62 attttgttgc ttagaaatca ccatgggtag 2070
64 <210> SEQ ID NO: 2
65 <211> LENGTH: 689
66 <212> TYPE: PRT
67 <213> ORGANISM: Homo sapiens
69 <400> SEQUENCE: 2
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71 1 5 10 15
72 Thr Ile Lys Thr Ile Lys Ala Glu Glu His Asn Lys Arg Gln Lys Glu
73 20 25 30
74 Arg Asn Val Thr Thr Gln Val Ser Val Asn Glu Ile Lys Gln Tyr Leu
75 35 40 45
76 Ser His Ile Leu Glu Gln Arg Thr Ser Ser Asn Val Ile Asn Lys Arg
77 50 55 60
78 Glu Asn Leu Leu Glu Lys Lys Lys Asn Gln Arg Lys Ile Arg Ile Lys
79 65 70 75 80
80 Gly Ile Gln Asn Lys Asp Ile Leu Lys Arg Asn Lys Asn His Leu Gln
81 85 90 95
82 Lys Gln Ala Glu Lys Asn Phe Thr Asp Glu Gly Asp Gln Leu Phe Lys
83 100 105 110
84 Met Gly Ile Lys Val Leu Gln Gln Ser Lys Ser Gln Lys Gln Lys Glu
85 115 120 125
86 Glu Ala Tyr Leu Leu Phe Ala Lys Ala Ala Asp Met Gly Asn Leu Lys
87 130 135 140
88 Ala Met Glu Lys Met Ala Asp Ala Leu Leu Phe Gly Asn Phe Gly Val
89 145 150 155 160
90 Gln Asn Ile Thr Ala Ala Ile Gln Leu Tyr Glu Ser Leu Ala Lys Glu
91 165 170 175
92 Gly Ser Cys Lys Ala Gln Asn Ala Leu Gly Phe Leu Ser Ser Tyr Gly
93 180 185 190
94 Ile Gly Met Glu Tyr Asp Gln Ala Lys Ala Leu Ile Tyr Tyr Thr Phe
95 195 200 205
96 Gly Ser Ala Gly Gly Asn Met Met Ser Gln Met Ile Leu Gly Tyr Arg
97 210 215 220
98 Tyr Leu Ser Gly Ile Asn Val Leu Gln Asn Cys Glu Val Ala Leu Ser
99 225 230 235 240
100 Tyr Tyr Lys Lys Val Ala Asp Tyr Ile Ala Asp Thr Phe Glu Lys Ser
101 245 250 255
102 Glu Gly Val Pro Val Glu Lys Val Arg Leu Thr Glu Arg Pro Glu Asn
103 260 265 270
104 Leu Ser Ser Asn Ser Glu Ile Leu Asp Trp Asp Ile Tyr Gln Tyr Tyr
105 275 280 285
106 Lys Phe Leu Ala Glu Arg Gly Asp Val Gln Ile Gln Val Ser Leu Gly

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156 Gly
159 <210> SEQ ID NO: 3
160 <211> LENGTH: 2067
161 <212> TYPE: DNA
162 <213> ORGANISM: Homo sapiens.
164 <400> SEQUENCE: 3
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166 atcaaaagcag aggaacataa taaaagacaa aaggaaaagaa atgtcaccac acagggtatca      120
167 gtgaacgaaa tcaaacataa tttatcacac atatttgaac aaagaacatc tagtaatgta      180
168 atcaataaaaa gagaaaatct cctggagaaa aagaagaatc aacgtaaaaat aagaataaaaa      240
169 ggaattcaaaa ataaagatat cttgaagaga aataagaatc atttacaaaa gcaagcagag      300
170 aaaaattttta cagatgaagg agaccagcta tttaagatgg gcatcaagggt tctccagcag      360
171 tctaaaagcc aaaaacaaaa agaagaagcc tacctacttt ttgccaaaagc agctgacatg      420
172 ggaaacttga aagctatgga gaaaatggct gacgctttgc tatttggaaa ttttggcgtg      480
173 caaaatataa cagcagctat ccaattatat gagtccttgg ctaaaagaagg atcatgtaaa      540
174 gcccaaacg cattaggatt tttgtcttct tatggaatag gaatggaata tgatcaagct      600
175 aaggcaactga tatattacac ctttgggaagt gctggaggaa acatgatgtc ccagatgatt      660
176 ttggggtaca gatatttgtc gggaatcaat gttctacaga atttgtgaagt tgccctaagt      720
177 tattacaaga aagtggcaga ttatattgct gacacatttg aaaaaagtga aggtgttcca      780
178 gtggaaaaaag tgagactaac ggaaagacct gaaaatctga gttctaacag tgagattttg      840
179 gattgggaca tataccaata ctataaattt ttggcagaaa gaggagatgt tcagatacaa      900
180 gtctctcttg gacaattaca tctaattggc aggaaggtc tagatcagga ttactacaaa      960
181 gcattacact acttctttaa ggcagcaaa ggcgggagtg caaatgccat ggcatttata      1020
182 ggaaagatgt atttagaggg gaatgctgcc gtgccgcaa ataacgctac tgccttcaag      1080
183 tacttttcca tggcagccag taagggcaat gcaatcggcc ttcattgggct tgggtcttctt      1140
184 tactttcatg gaaaaggagt tcccctgaat tatgccraag cacttaaata ctttcagaaa      1200
185 gctgcggaaa aagggtggcc cgacgcacag ttccagttag gcttcatgta ctactctggc      1260
186 tctggaatat ggaaggatta taaacttgcc ttcaaattt tttacctggc atctcagagt      1320
187 gggcagcccc tcgccattta ttatctggcc aagatgtatg caacaggaac aggagtagta      1380
188 agatcatgca gaactgctgt ggagctttat aaagggtgtc gtgaactagg ccaactgggct      1440
189 gagaaattcc tgacagctta ctttgcctat aaggatgggtg atatagattc ttctcttggt      1500
190 cagtatgcac tgcttgacaga aatgggggtat gaagtagctc aaagcaattc agcattcatt      1560
191 ttggaatcta aaaaggctaa cattcttgaa aaagagaaga tgtatccaat ggcgcttctc      1620
192 ctatggaatc gagctgccat tcaaggcaat gcatttgcta gagtaaaaaat tggagattac      1680
193 cattactatg gctatgggac taagaaagac tatcaaacag cagccacaca ctacagcatt      1740
194 gcagccaaca aataccacaa cgcgcaagcc atgttcaatc tggcttatat gtatgaacac      1800
195 ggcttaggca tcacaaaagga cattcacttg gccagaagat tgtacgacat ggctgctcaa      1860
196 acgagtccag atgcccacat acctgtgctc tttgccgtca tgaaactgga aactacgcat      1920
197 ttgtctcggg atatctgttt ttttaatttc acaacgagat ggaactggct gaaactggac      1980
198 aacaccattg gaccacactg ggacttattt gtgattggcc tcattgttcc tgggctgatt      2040
199 ttgttgctta gaaatcacca tgggtag      2067
201 <210> SEQ ID NO: 4
202 <211> LENGTH: 688
203 <212> TYPE: PRT
204 <213> ORGANISM: Homo sapiens
206 <400> SEQUENCE: 4
207 Met Lys Pro Leu Ser Leu Leu Ile Glu Ile Leu Ile Ile Leu Gly Val
208 1 5 10 15
209 Thr Ile Lys Thr Ile Lys Ala Glu Glu His Asn Lys Arg Gln Lys Glu

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210		20		25		30
211	Arg Asn Val Thr Thr Gln Val Ser Val Asn Glu Ile Lys Gln Tyr Leu					
212		35		40		45
213	Ser His Ile Leu Glu Gln Arg Thr Ser Ser Asn Val Ile Asn Lys Arg					
214		50		55		60
215	Glu Asn Leu Leu Glu Lys Lys Lys Asn Gln Arg Lys Ile Arg Ile Lys					
216	65		70		75	80
217	Gly Ile Gln Asn Lys Asp Ile Leu Lys Arg Asn Lys Asn His Leu Gln					
218		85		90		95
219	Lys Gln Ala Glu Lys Asn Phe Thr Asp Glu Gly Asp Gln Leu Phe Lys					
220		100		105		110
221	Met Gly Ile Lys Val Leu Gln Gln Ser Lys Ser Gln Lys Gln Lys Glu					
222		115		120		125
223	Glu Ala Tyr Leu Leu Phe Ala Lys Ala Ala Asp Met Gly Asn Leu Lys					
224		130		135		140
225	Ala Met Glu Lys Met Ala Asp Ala Leu Leu Phe Gly Asn Phe Gly Val					
226	145		150		155	160
227	Gln Asn Ile Thr Ala Ala Ile Gln Leu Tyr Glu Ser Leu Ala Lys Glu					
228		165		170		175
229	Gly Ser Cys Lys Ala Gln Asn Ala Leu Gly Phe Leu Ser Ser Tyr Gly					
230		180		185		190
231	Ile Gly Met Glu Tyr Asp Gln Ala Lys Ala Leu Ile Tyr Tyr Thr Phe					
232		195		200		205
233	Gly Ser Ala Gly Gly Asn Met Met Ser Gln Met Ile Leu Gly Tyr Arg					
234		210		215		220
235	Tyr Leu Ser Gly Ile Asn Val Leu Gln Asn Cys Glu Val Ala Leu Ser					
236	225		230		235	240
237	Tyr Tyr Lys Lys Val Ala Asp Tyr Ile Ala Asp Thr Phe Glu Lys Ser					
238		245		250		255
239	Glu Gly Val Pro Val Glu Lys Val Arg Leu Thr Glu Arg Pro Glu Asn					
240		260		265		270
241	Leu Ser Ser Asn Ser Glu Ile Leu Asp Trp Asp Ile Tyr Gln Tyr Tyr					
242		275		280		285
243	Lys Phe Leu Ala Glu Arg Gly Asp Val Gln Ile Gln Val Ser Leu Gly					
244		290		295		300
245	Gln Leu His Leu Ile Gly Arg Lys Gly Leu Asp Gln Asp Tyr Tyr Lys					
246	305		310		315	320
247	Ala Leu His Tyr Phe Leu Lys Ala Ala Lys Ala Gly Ser Ala Asn Ala					
248		325		330		335
249	Met Ala Phe Ile Gly Lys Met Tyr Leu Glu Gly Asn Ala Ala Val Pro					
250		340		345		350
251	Gln Asn Asn Ala Thr Ala Phe Lys Tyr Phe Ser Met Ala Ala Ser Lys					
252		355		360		365
253	Gly Asn Ala Ile Gly Leu His Gly Leu Gly Leu Leu Tyr Phe His Gly					
254		370		375		380
255	Lys Gly Val Pro Leu Asn Tyr Ala Glu Ala Leu Lys Tyr Phe Gln Lys					
256	385		390		395	400
257	Ala Ala Glu Lys Gly Trp Pro Asp Ala Gln Phe Gln Leu Gly Phe Met					
258		405		410		415

## VERIFICATION SUMMARY

DATE: 05/29/2001

PATENT APPLICATION: US/09/714,882

TIME: 11:46:14

Input Set : A:\Seqlist.txt

Output Set: C:\CRF3\05292001\I714882.raw

L:15 M:270 C: Current Application Number differs, Replaced Current Application No

L:15 M:271 C: Current Filing Date differs, Replaced Current Filing Date